

RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number:	10/622,845
Source:	TFWO.
Date Processed by STIC:	12/23/03
~ 4	

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.
PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

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- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 703-308-4212; FAX: 703-308-4221 <u>Effective 12/13/03</u>: TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 4.1 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

http://www.uspto.gov/web/offices/pac/checker/chkr41note.htm

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail. Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom. Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

- 1. EFS-Bio (http://www.uspto.gov/ebc/efs/downloads/documents.htm, EFS Submission User Manual ePAVE)
- 2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
- Hand Carry directly to (EFFECTIVE 12/01/03):
 U.S. Patent and Trademark Office, Box Sequence, Customer Window, Lobby, Room 1B03, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202
- Federal Express, United Parcel Service, or other delivery service to: U.S. Patent and Trademark Office, Box Sequence, Room 1B03-Mailroom, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202

Revised 10/08/03



IFWO

RAW SEQUENCE LISTING DATE: 12/30/2003 PATENT APPLICATION: US/10/622,893 TIME: 09:13:52

Input Set: N:\Crf4\12232003\J622893.raw
Output Set: N:\CRF4\12302003\J622893.raw

```
Collegied Diskette Mesded
              1 <110> APPLICANT: Yuan, Chong-Sheng
                                                                                                                                                                 Dices Prov. Comply
              2
                                 Datta, Abhijit
                                 Wang, Yuping
              4 <120> TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR
                                  DETERMINATION OF GLYCATED PROTEINS
              6 <130> FILE REFERENCE: 466992001300
                                                                                                           - Invalid Response Fus (2137.

Asp T.
              7 <140> CURRENT APPLICATION NUMBER: US/10/622,893
C--> 8 <141> CURRENT FILING DATE: 2003-07-17
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            16 <220> FEATURE:
            17 <223> OTHER INFORMATION: / peptide
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            19
            20
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            23 <211> LENGTH: 6
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            26 <220> FEATURE:
            27 <223> OTHER INFORMATION: FAD cofactor-binding consensus sequence
            28 <220> FEATURE:
            29 <221> NAME/KEY: VARIANT
            30 <222> LOCATION: 2, 4, 5
            31 <223> OTHER INFORMATION: Xaa = Any Amino Acid
            32 <400> SEQUENCE: 2
W--> 33
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                                                                                                                   - Invalid Response, ports

- Invalid Response, p
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            38 <212> TYPE: PRT
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            40 <220> FEATURE:
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            44
                                                                          5
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            45
                                                                20
                                                                                                                  25
```

Input Set : N:\Crf4\12232003\J622893.raw
Output Set: N:\CRF4\12302003\J622893.raw

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49 50	Asp	Val 50	Asn	Lys	Val	Ile	Ser 55		Gly	Gln	Tyr	Ser 60		Asn	Lys	Asp
51 52	Glu 65		Glu	Val	Asn	Glu 70		Leu	Ala	Glu	Glu 75	Ala	Phe	Asn	Gly	Trp 80
53 54	Lys	Asn	Asp	Pro	Leu 85	Phe	Lys	Pro	Tyr	Tyr 90	His	Asp	Thr	Gly	Leu 95	Leu
55 56			Ala	100				_	105				-	110		
57 58	-		Gly 115		_			120					125			
59 60		130	Lys				135	_				140				
61 62	145	-	Gly			150					155					160
63 64	Asn	Ala	Leu	Val	Ala 165	Ala	Ala	Arg	Glu	Ala 170	Gln	Arg	Met	Gly	Val 175	Lys
65 66			Thr	180				_	185					190		
67 68			Asp 195			_		200					205			
69 70		210	Arg				215					220				
71 72	Asp 225	Phe	Lys	Asn	Gln	Leu 230	Arg	Pro	Thr	Ala	Trp 235	Thr	Leu	Val	His	Ile 240
73 74			Lys		245					250					255	
75 76			Ile	260	_				265					270		
77 78			Lys 275					280					285			
79 80		290	Asp	_			295					300				
81 82	305		Glu			310	_				315					320
83 84			Leu		325					330					335	
85 86		_	Thr	340					345					350		
87 88			Leu 355			_	_	360					365			
89 90		370	Ser				375					380				
91 92	385		Lys			390					395					400
93 94	Asn	Arg	Asn	Trp	Arg 405	Asp	Thr	Leu	Gly	Arg 410	Phe	Gly	Gly	Pro	Asn 415	Arg
95	Val	Met	Asp	Phe	His	Asp	Val	Lys	Glu	Trp	Thr	Asn	Val	Gln	Tyr	Arg

Input Set : N:\Crf4\12232003\J622893.raw
Output Set: N:\CRF4\12302003\J622893.raw

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430
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96
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97
98
                 435
                                         , s ame error
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101 <211> LENGTH: 17
102 <212> TYPE: PRT
103 <213> ORGANISM: Artificial Sequence
104 <220> FEATURE:
105 <223> OTHER INFORMATION (
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106 <400> SEQUENCE: 4
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107
108
          Gly
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112 <211> LENGTH: 472
113 <212> TYPE: PRT
114 <213> ORGANISM: Artificial Sequence
115 <220> FEATURE:
116 <223> OTHER INFORMATION: chimeric protein
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120
121
                      20
                                          25
          Ala Leu His Leu Ala Arg Arg Gly Tyr Thr Asn Val Thr Val Leu Asp
122
123
          Pro Tyr Pro Val Pro Ser Ala Ile Ser Ala Gly Asn Asp Val Asn Lys
124
125
                                  55
          Val Ile Ser Ser Gly Gln Tyr Ser Asn Asn Lys Asp Glu Ile Glu Val
126
                              70
                                                   75
127
          Asn Glu Ile Leu Ala Glu Glu Ala Phe Asn Gly Trp Lys Asn Asp Pro
128
129
          Leu Phe Lys Pro Tyr Tyr His Asp Thr Gly Leu Leu Met Ser Ala Cys
130
131
                      100
          Ser Gln Glu Gly Leu Asp Arg Leu Gly Val Arg Val Arg Pro Gly Glu
132
133
                                      120
          Asp Pro Asn Leu Val Glu Leu Thr Arg Pro Glu Gln Phe Arg Lys Leu
134
                                                       140
                                  135
135
          Ala Pro Glu Gly Val Leu Gln Gly Asp Phe Pro Gly Trp Lys Gly Tyr
136
                              150
                                                   155
137
          Phe Ala Arg Ser Gly Ala Gly Trp Ala His Ala Arg Asn Ala Leu Val
138
139
                          165
                                              170
          Ala Ala Arg Glu Ala Gln Arg Met Gly Val Lys Phe Val Thr Gly
140
141
                                          185
          Thr Pro Gln Gly Arg Val Val Thr Leu Ile Phe Glu Asn Asn Asp Val
142
                                      200
143
          Lys Gly Ala Val Thr Gly Asp Gly Lys Ile Trp Arg Ala Glu Arg Thr
144
145
          Phe Leu Cys Ala Gly Ala Ser Ala Gly Gln Phe Leu Asp Phe Lys Asn
146
```

Input Set: N:\Crf4\12232003\J622893.raw
Output Set: N:\CRF4\12302003\J622893.raw

```
240
                              230
                                                   235
147
          225
         Gln Leu Arg Pro Thr Ala Trp Thr Leu Val His Ile Ala Leu Lys Pro
148
                                               250
149
                          245
         Glu Glu Arg Ala Leu Tyr Lys Asn Ile Pro Val Ile Phe Asn Ile Glu
150
                                                               270
151
                      260
                                          265
         Arg Gly Phe Phe Glu Pro Asp Glu Glu Arg Gly Glu Ile Lys Ile
152
153
                  275
                                      280
          Cys Asp Glu His Pro Gly Tyr Thr Asn Met Val Gln Ser Ala Asp Gly
154
155
                                  295
                                                       300
          Thr Met Met Ser Ile Pro Phe Glu Lys Thr Gln Ile Pro Lys Glu Ala
156
                              310
                                                   315
157
          Glu Thr Arg Val Arg Ala Leu Leu Lys Glu Thr Met Pro Gln Leu Ala
158
                                               330
159
                          325
          Asp Arg Pro Phe Ser Phe Ala Arg Ile Cys Trp Cys Ala Asp Thr Ala
160
                                           345
                      340
161
          Asn Arg Glu Phe Leu Ile Asp Arg His Pro Gln Tyr His Ser Leu Val
162
163
                                       360
          Leu Gly Cys Gly Ala Ser Gly Arg Gly Phe Lys Tyr Leu Pro Ser Ile
164
                                  375
                                                       380
165
          Gly Asn Leu Ile Val Asp Ala Met Glu Gly Lys Val Pro Gln Lys Ile
166
                              390
                                                   395
167
          His Glu Leu Ile Lys Trp Asn Pro Asp Ile Ala Ala Asn Arg Asn Trp
168
                                               410
169
                          405
          Arg Asp Thr Leu Gly Arg Phe Gly Gly Pro Asn Arg Val Met Asp Phe
170
                                           425
171
                      420
          His Asp Val Lys Glu Trp Thr Asn Val Gln Tyr Arg Asp Ile Ser Lys
172
                                       440
                                                           445
173
          Leu Lys Gly Glu Leu Glu Gly Leu Pro Ile Pro Asn Pro Leu Leu Arg
174
                                   455
175
                                         Nucleofide
          Thr Gly His His His His His
176
177
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180 <211> LENGTH: 1419
181 <212> TYPE: DNA
182 <213> ORGANISM: Artificial Sequence
183 <220> FEATURE:
184 <223> OTHER INFORMATION: /nuc/sequence encoding a chimeric protein
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186
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187
          tataccaacg ttaccgtgct ggacccctat cctgtcccta gcgccatctc cgccggaaac 180
188
          gacgtgaaca aagtcattag cagtggccaa tattcgaata acaaagacga aatcgaagtg 240
189
          aatgagatet tggeggaaga ggegtttaac ggttggaaga acgaeceget tttcaaaceg 300
190
          tattatcatg atacgggcct gctgatgtct gcttgctcgc aggagggcct ggatcgcctg 360
191
          ggcgtccggg tacgtccggg cgaggatcct aatctggtgg aacttacccg cccggagcaa 420
192
          tttcgtaaac tggccccgga aggcgtgttg caaggtgatt ttccgggttg gaaagggtac 480
193
          tttgcgcgtt ccggcgctgg ctgggcacat gcaaggaatg ccttagtggc agcagcacgc 540
194
          qaaqcacaqc gcatgggtgt aaaatttgtt actggcaccc cgcagggtcg tgtagtcacg 600
195
          ttaatctttg aaaataacga tgtaaaaggt gccgttacgg gcgatggcaa aatttggaga 660
196
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Input Set : N:\Crf4\12232003\J622893.raw
Output Set: N:\CRF4\12302003\J622893.raw

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    197
              caacttcgac caaccgcttg gaccctggta cacattgcgt taaaaccgga agaacgtgcg 780
    198
              ttgtacaaaa atataccggt tatctttaac atcgaacggg ggtttttctt tgaacccgat 840
    199
              qaqqaqcqcq qtqagattaa aatatgcgat gaacacccgg gctacacaaa tatggtccag 900
    200
              agtgcagacg gcacgatgat gagcattccg ttcgaaaaaa cccagattcc aaaagaagcc 960
    201
              gaaacgcgcg ttcgggccct gctgaaagag acaatgcccc agctggcaga ccgtccattc 1020
    202
              agettegeae geatttgetg gtgtgeegat aeegegaate gegaatteet gatagatega 1080
    203
              catccgcagt accacagtct tgtgttgggc tgtggtgcga.gcggaagagg gtttaaatat 1140
    204
              ctqccttcta ttgggaatct cattgttgac gcgatggaag gtaaagtgcc gcaaaaaatt 1200
    205
              cacqaattaa tcaagtggaa cccggacatt gcggcgaacc gtaactggcg tgatactctg 1260
    206
              gggcgttttg gcggtccaaa tcgtgtgatg gattttcatg atgtgaagga atggaccaat 1320
    207
              gttcagtatc gtgatatttc caagctgaaa ggagagttgg aaggtaagcc aatccctaac 1380
    208
              ccgttactgc gcacaggcca tcaccatcat catcattaa
    209
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                                        Same erect
    212 <211> LENGTH: 39
    213 <212> TYPE: PRT
    214 <213> ORGANISM: Artificial Sequence
    215 <220> FEATURE:
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    216 <223> OTHER INFORMATION
    217 <220> FEATURE:
    218 <221> NAME/KEY: VARIANT
    219 <222> LOCATION: 12
    220 <223> OTHER INFORMATION: Xaa = C or T
    221 <400> SEQUENCE: 7
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W--> 222
                                                10
    223
              224
                                              Same erver
                                            25
                         20
    225
              Gly Gly Gly Gly Gly Gly
    226
    227
                      35
    229 <210> SEO ID NO: 8
    230 <211> LENGTH: 39
    231 <212> TYPE: PRT
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    233 <220> FEATURE:
    234 <223> OTHER INFORMATION;
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    237
              238
                                            25
    239
                         20
              Gly Gly Gly Gly Gly Gly
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                      35
    241
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    244 <211> LENGTH: 39
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    246 <213> ORGANISM: Artificial Sequence
    247 <220> FEATURE:
    248 <223> OTHER INFORMATION: peptide
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RAW SEQUENCE LISTING ERROR SUMMARY PATENT APPLICATION: US/10/622,893

DATE: 12/30/2003 TIME: 09:13:53

Input Set : N:\Crf4\12232003\J622893.raw
Output Set: N:\CRF4\12302003\J622893.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:2; Xaa Pos. 2,4,5
Seq#:7; Xaa Pos. 12

VERIFICATION SUMMARY

PATENT APPLICATION: US/10/622,893

DATE: 12/30/2003 TIME: 09:13:53

Input Set : N:\Crf4\12232003\J622893.raw
Output Set: N:\CRF4\12302003\J622893.raw

L:8 M:271 C: Current Filing Date differs, Replaced Current Filing Date L:33 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:0 L:222 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7 after pos.:0